

0420  
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## RAW SEQUENCE LISTING

DATE: 02/19/2002

PATENT APPLICATION: US/10/066,960

TIME: 12:40:15

Input Set : N:\Crf3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

## SEQUENCE LISTING

2 (1) GENERAL INFORMATION:  
 3 (i) APPLICANT: PARMA, et al.  
 4 (ii) TITLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID LIGANDS  
 5 TO LECTINS  
 6 (iii) NUMBER OF SEQUENCES: 390  
 7 (iv) CORRESPONDENCE ADDRESS:  
 8 (A) ADDRESSEE: Swanson & Bratschun, L.L.C.  
 9 (B) STREET: 8400 E. Prentice Avenue, Suite 200  
 10 (C) CITY: Englewood  
 11 (D) STATE: Colorado  
 12 (E) COUNTRY: USA  
 13 (F) ZIP: 80111  
 14 (v) COMPUTER READABLE FORM:  
 15 (A) MEDIUM TYPE: Diskette, 3 1/2 diskette, 1.44 MB  
 16 (B) COMPUTER: IBM pc compatible  
 17 (C) OPERATING SYSTEM: MS-DOS  
 18 (D) SOFTWARE: WordPerfect 6.0  
 19 (vi) CURRENT APPLICATION DATA:  
 C--> 20 (A) APPLICATION NUMBER: US/10/066,960  
 C--> 21 (B) FILING DATE: 04-Feb-2002  
 26 (C) CLASSIFICATION:  
 39 (vii) PRIOR APPLICATION DATA:  
 23 (A) APPLICATION NUMBER: 08/952,793  
 24 (B) FILING DATE: 1999-DEC-03  
 28 (A) APPLICATION NUMBER: PCT/US96/09455  
 29 (B) FILING DATE: 05-JUNE-1995  
 31 (A) APPLICATION NUMBER: 08/479,724  
 32 (B) FILING DATE: 07-JUNE-1995  
 34 (A) APPLICATION NUMBER: 08/472,256  
 35 (B) FILING DATE: 07-JUNE-1995  
 37 (A) APPLICATION NUMBER: 08/472,255  
 38 (B) FILING DATE: 07-JUNE-1995  
 40 (A) APPLICATION NUMBER: 08/477,829  
 41 (B) FILING DATE: 07-JUNE-1995  
 42 (viii) ATTORNEY/AGENT INFORMATION:  
 43 (A) NAME: Barry J. Swanson  
 44 (B) REGISTRATION NUMBER: 33,215  
 45 (C) REFERENCE/DOCKET NUMBER: NEX40C/PCT  
 46 (ix) TELECOMMUNICATION INFORMATION:  
 47 (A) TELEPHONE: (303) 793-3333  
 48 (B) TELEFAX: (303) 793-3433  
 50 (2) INFORMATION FOR SEQ ID NO: 1:

ENTERED

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Input Set : N:\Crif3\RULE60\10066960.txt

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51      (i) SEQUENCE CHARACTERISTICS:
52          (A) LENGTH: 98 base pairs
53          (B) TYPE: nucleic acid
54          (C) STRANDEDNESS: single
55          (D) TOPOLOGY: linear
W--> 56      (ii) MOLECULE TYPE: RNA
57      (ix) FEATURE:
58          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
59      (ix) FEATURE:
60          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
61      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
62      GGGAAAAGCG AAUCAUACAC AAGANNNNNNN NNNNNNNNNNN NNNNNNNNNNN 50
63      NNNNNNNNNNN NNNNNNNNNNN NNNNGCUCCG CCAGAGACCA ACCGAGAA 98
65 (2) INFORMATION FOR SEQ ID NO: 2:
66      (i) SEQUENCE CHARACTERISTICS:
67          (A) LENGTH: 41 base pairs
68          (B) TYPE: nucleic acid
69          (C) STRANDEDNESS: single
70          (D) TOPOLOGY: linear
W--> 71      (ii) MOLECULE TYPE: RNA
72      (ix) FEATURE:
73          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
74      (ix) FEATURE:
75          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
76      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
77      UAAUACGACU CACUAUAGGG AAAAGCGAAU CAUACACAAG A 41
79 (2) INFORMATION FOR SEQ ID NO: 3:
80      (i) SEQUENCE CHARACTERISTICS:
81          (A) LENGTH: 24 base pairs
82          (B) TYPE: nucleic acid
83          (C) STRANDEDNESS: single
84          (D) TOPOLOGY: linear
W--> 85      (ii) MOLECULE TYPE: RNA
86      (ix) FEATURE:
87          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
88      (ix) FEATURE:
89          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
90      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
91      UUCUCGGUUG GUCUCUGGCG GAGC 24
93 (2) INFORMATION FOR SEQ ID NO: 4:
94      (i) SEQUENCE CHARACTERISTICS:
95          (A) LENGTH: 96 base pairs
96          (B) TYPE: nucleic acid
97          (C) STRANDEDNESS: single
98          (D) TOPOLOGY: linear
W--> 99      (ii) MOLECULE TYPE: RNA
100     (ix) FEATURE:
101         (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
102     (ix) FEATURE:

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103      (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
104      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
105      GGGAAAAGCG AAUCAUACAC AAGAAUGGUU GGCCUGGGCG CAGGCUUCGA      50
106      AGACUCGGCG GGAACGGGAA UGGCUCGCC AGAGACCAAC CGAGAA      96
108 (2) INFORMATION FOR SEQ ID NO: 5:
109      (i) SEQUENCE CHARACTERISTICS:
110          (A) LENGTH: 98 base pairs
111          (B) TYPE: nucleic acid
112          (C) STRANDEDNESS: single
113          (D) TOPOLOGY: linear
W--> 114      (ii) MOLECULE TYPE: RNA
115      (ix) FEATURE:
116          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
117      (ix) FEATURE:
118          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
119      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
120      GGGAAAAGCG AAUCAUACAC AAGACAGGCA CUGAAAACUC GCGGGAACG      50
121      AAAGUAGUGC CGACUCAGAC GCGUGCUCG CCAGAGACCA ACCGAGAA      98
123 (2) INFORMATION FOR SEQ ID NO: 6:
124      (i) SEQUENCE CHARACTERISTICS:
125          (A) LENGTH: 91 base pairs
126          (B) TYPE: nucleic acid
127          (C) STRANDEDNESS: single
128          (D) TOPOLOGY: linear
W--> 129      (ii) MOLECULE TYPE: RNA
130      (ix) FEATURE:
131          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
132      (ix) FEATURE:
133          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
134      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
135      GGGAAAAGCG AAUCAUACAC AAGAAGUCUG GCCAAAGACU CGGCGGGAAC      50
136      GUAAAACGGC CAGAAUUGCU CCGCCAGAGA CCAACCGAGA A      91
138 (2) INFORMATION FOR SEQ ID NO: 7:
139      (i) SEQUENCE CHARACTERISTICS:
140          (A) LENGTH: 94 base pairs
141          (B) TYPE: nucleic acid
142          (C) STRANDEDNESS: single
143          (D) TOPOLOGY: linear
W--> 144      (ii) MOLECULE TYPE: RNA
145      (ix) FEATURE:
146          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
147      (ix) FEATURE:
148          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
149      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
150      GGGAAAAGCG AAUCAUACAC AAGAGUAGGA GGUUCCAUCA CCAGGACUCG      50
151      GCGGGAACG AAGGUGAUGS GCUCCGCCAG AGACCAACCG AGAA      94
153 (2) INFORMATION FOR SEQ ID NO: 8:
154      (i) SEQUENCE CHARACTERISTICS:
155          (A) LENGTH: 95 base pairs

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156         (B) TYPE: nucleic acid
157         (C) STRANDEDNESS: single
158         (D) TOPOLOGY: linear
W--> 159     (ii) MOLECULE TYPE: RNA
160         (ix) FEATURE:
161             (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
162         (ix) FEATURE:
163             (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
164         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
165             GGGAAAAGCG AAUCAUACAC AAGAACAAGG AUCGAUGGCG AGCCGGGGAG      50
166             GGCUCGGCGG GAACGAAUUC UGCUCGCCCA GAGACCAACC GAGAA          95
168 (2) INFORMATION FOR SEQ ID NO: 9:
169     (i) SEQUENCE CHARACTERISTICS:
170         (A) LENGTH: 97 base pairs
171         (B) TYPE: nucleic acid
172         (C) STRANDEDNESS: single
173         (D) TOPOLOGY: linear
W--> 174     (ii) MOLECULE TYPE: RNA
175         (ix) FEATURE:
176             (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
177         (ix) FEATURE:
178             (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
179         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
180             GGGAAAAGCG AAUCAUACAC AAGAUUGGGC AGGCAGAGCG AGACCGGGGG      50
181             CUCGGCGGGA ACGGAACAGG AAUGCUCGCG CAGAGACCAA CCGAGAA          97
183 (2) INFORMATION FOR SEQ ID NO: 10:
184     (i) SEQUENCE CHARACTERISTICS:
185         (A) LENGTH: 97 base pairs
186         (B) TYPE: nucleic acid
187         (C) STRANDEDNESS: single
188         (D) TOPOLOGY: linear
W--> 189     (ii) MOLECULE TYPE: RNA
190         (ix) FEATURE:
191             (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
192         (ix) FEATURE:
193             (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
194         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
195             GGGAAAAGCG AAUCAUACAC AAGAAAGGGA UGGGAUUGGG ACGAGCGGCC      50
196             AAGACUCGGC GGGAACGAAG GGUGCUCGCG CAGAGACCAA CCGAGAA          97
198 (2) INFORMATION FOR SEQ ID NO: 11:
199     (i) SEQUENCE CHARACTERISTICS:
200         (A) LENGTH: 96 base pairs
201         (B) TYPE: nucleic acid
202         (C) STRANDEDNESS: single
203         (D) TOPOLOGY: linear
W--> 204     (ii) MOLECULE TYPE: RNA
205         (ix) FEATURE:
206             (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
207         (ix) FEATURE:

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208          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
209      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
210          GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGAAA GUGUCAUGGU      50
211          AGCAAGUCCA AUGGUGGACU CUGCUCGCC AGAGACCAAC CGAGAA      96
213 (2) INFORMATION FOR SEQ ID NO: 12:
214      (i) SEQUENCE CHARACTERISTICS:
215          (A) LENGTH: 98 base pairs
216          (B) TYPE: nucleic acid
217          (C) STRANDEDNESS: single
218          (D) TOPOLOGY: linear
W--> 219      (ii) MOLECULE TYPE: RNA
220      (ix) FEATURE:
221          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
222      (ix) FEATURE:
223          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
224      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
225          GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGUGA AGUGGGUAGG      50
226          UAGCUGAAGA CGGUCUGGGC GCCAGCUCCG CCAGAGACCA ACCGAGAA      98
228 (2) INFORMATION FOR SEQ ID NO: 13:
229      (i) SEQUENCE CHARACTERISTICS:
230          (A) LENGTH: 99 base pairs
231          (B) TYPE: nucleic acid
232          (C) STRANDEDNESS: single
233          (D) TOPOLOGY: linear
W--> 234      (ii) MOLECULE TYPE: RNA
235      (ix) FEATURE:
236          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
237      (ix) FEATURE:
238          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
239      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
240          GGGAAAAGCG AAUCAUACAC AAGAAAGGGA UGGGAUUGGG ACGAGCGGCC      50
241          AAGACUCGGC GGGAACGAAG GUUCCGCUCC GCCAGAGACC AACCGAGAA      99
243 (2) INFORMATION FOR SEQ ID NO: 14:
244      (i) SEQUENCE CHARACTERISTICS:
245          (A) LENGTH: 98 base pairs
246          (B) TYPE: nucleic acid
247          (C) STRANDEDNESS: single
248          (D) TOPOLOGY: linear
W--> 249      (ii) MOLECULE TYPE: RNA
250      (ix) FEATURE:
251          (D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
252      (ix) FEATURE:
253          (D) OTHER INFORMATION: All U's are 2'-NH2 uracil
254      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
255          GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGAAG UGUGUGAGUA      50
256          ACGAUCACUU GGUACUAAAA GCCCGCUCCG CCAGAGACCA ACCGAGAA      98
258 (2) INFORMATION FOR SEQ ID NO: 15:
259      (i) SEQUENCE CHARACTERISTICS:
260          (A) LENGTH: 100 base pairs

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DATE: 02/19/2002

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Input Set : N:\CrF3\RULE60\10066960.txt

Output Set: N:\CRF3\02192002\J066960.raw

L:20 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]  
L:21 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]  
L:56 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:56 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=1  
L:71 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:71 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=2  
L:85 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:85 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3  
L:99 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:99 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4  
L:114 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:114 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=5  
L:129 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:129 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=6  
L:144 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:144 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=7  
L:159 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:159 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=8  
L:174 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:174 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=9  
L:189 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:189 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=10  
L:204 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:204 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=11  
L:219 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:219 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=12  
L:234 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:234 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=13  
L:249 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:249 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=14  
L:264 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:264 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=15  
L:279 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:279 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=16  
L:295 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:295 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=17  
L:310 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
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L:325 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
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L:340 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:340 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=20  
L:355 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:355 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=21  
L:370 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:370 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=22  
L:385 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:385 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=23

## VERIFICATION SUMMARY

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Input Set : N:\Crif3\RULE60\10066960.txt

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L:400 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:400 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=24  
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L:415 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=25  
L:430 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
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L:445 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:445 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=27  
L:460 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:460 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=28  
L:475 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:475 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=29  
L:490 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:490 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=30  
L:505 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:505 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=31  
L:520 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:520 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=32  
L:535 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:535 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=33  
L:550 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:550 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=34  
L:565 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:565 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=35  
L:580 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:580 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=36  
L:595 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:595 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=37  
L:610 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:610 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=38  
L:625 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:625 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=39  
L:640 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:640 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=40  
L:655 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:655 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=41  
L:670 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:670 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=42  
L:685 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:685 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=43  
L:700 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:700 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=44  
L:715 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:715 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=45  
L:730 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:730 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=46  
L:745 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]  
L:745 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=47  
L:760 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]

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Input Set : N:\Crf3\RULE60\10066960.txt

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L:760 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=48  
L:775 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=49  
L:790 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=50